



RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 10/712,533
Source: OIDE
Date Processed by STIC: 8-10-04

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
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1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
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IFWO

RAW SEQUENCE LISTING

DATE: 08/10/2004

PATENT APPLICATION: US/10/712,533

TIME: 11:50:29

Input Set : A:\PTO.LM.txt

Output Set: N:\CRF4\08102004\J712533.raw

5 <110> APPLICANT: Institut Pasteur

9 <120> TITLE OF INVENTION: ANTIGENES DE PLASMODIUM FALCIPARUM ET LEURS APPLICATIONS
VACCINALES ET DIAGNOSTIQUES

10

14 <130> FILE REFERENCE: 000466-0035

C--> 18 <140> CURRENT APPLICATION NUMBER: US/10/712,533

C--> 18 <141> CURRENT FILING DATE: 2003-11-14

18 <150> PRIOR APPLICATION NUMBER: CA 2,345,206

20 <151> PRIOR FILING DATE: 2001-05-16

24 <150> PRIOR APPLICATION NUMBER: CA 2,346,968

26 <151> PRIOR FILING DATE: 2001-05-23

30 <160> NUMBER OF SEQ ID NOS: 18

34 <170> SOFTWARE: PatentIn version 3.1

38 <210> SEQ ID NO: 1

40 <211> LENGTH: 192

42 <212> TYPE: DNA

44 <213> ORGANISM: Plasmodium falciparum

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51 aatgtttataa aaagtataa taaaaatgtt ataaaaagtg ataataaaaa tgattataaa 120

53 aagtgtataa aaaatgttat aaaaagtgtat aataaaaaatg ttataaaaaag tgataataaa 180

55 aatgtggaat tc 192

58 <210> SEQ ID NO: 2

60 <211> LENGTH: 351

62 <212> TYPE: DNA

64 <213> ORGANISM: Plasmodium falciparum

68 <400> SEQUENCE: 2

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71 agtagagatg ataataatag tctctctaag agcgtagata atcaaagtaa tgttttaaat 120

73 agagaagatc ctattgcttc tgaaactgaa gttgttaagt aacctgagga ttcaagtagg 180

75 ataatgacta cagaagttcc aagtactact gtaaaacccc ctgatgaaaa acgatctgaa 240

77 gaagtaggag aaaaagaagc taaagaaatt aaagtagaac ctgttgtagc aagagccatt 300

79 ggagaaccaa tggaaaattc tgtgagcgta cagtccectc ctaaggaatt c 351

82 <210> SEQ ID NO: 3

84 <211> LENGTH: 64

86 <212> TYPE: PRT

88 <213> ORGANISM: Plasmodium falciparum

92 <400> SEQUENCE: 3

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95 1 5 10 15

98 Asp Lys Glu Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys

99 20 25 30

102 Ser Asp Asn Lys Asn Asp Tyr Lys Lys Cys Asn Lys Asn Val Ile Lys

103 35 40 45

*please use english
for US. applications*

**Does Not Comply
Corrected Diskette Needed**

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Input Set : A:\PTO.LM.txt

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127      20                      25                      30
130 Asp Asn Gln Ser Asn Val Leu Asn Arg Glu Asp Pro Ile Ala Ser Glu
131      35                      40                      45
134 Thr Glu Val Val Ser Glu Pro Glu Asp Ser Ser Arg Ile Met Thr Thr
135      50                      55                      60
138 Glu Val Pro Ser Thr Thr Val Lys Pro Pro Asp Glu Lys Arg Ser Glu
139 65                      70                      75                      80
142 Glu Val Gly Glu Lys Glu Ala Lys Glu Ile Lys Val Glu Pro Val Val
143      85                      90                      95
146 Pro Arg Ala Ile Gly Glu Pro Met Glu Asn Ser Val Ser Val Gln Ser
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150 Pro Pro Lys Glu Phe
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188 <211> LENGTH: 6
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204 <211> LENGTH: 7
206 <212> TYPE: PRT
208 <213> ORGANISM: Plasmodium falciparum
212 <400> SEQUENCE: 8

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Input Set : A:\PTO.LM.txt

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232 <222> LOCATION: (1)..(1524)
234 <223> OTHER INFORMATION:
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240 Met Lys Thr Thr Lys Glu Asn Asp Asn Asn Asn Ile Val His Tyr Val
241 1          5          10          15
243 gat tgg ata aac cag att ttt aaa aag aat tct tta caa tgt gat tta      96
244 Asp Trp Ile Asn Gln Ile Phe Lys Lys Asn Ser Leu Gln Cys Asp Leu
245          20          25          30
247 tat ttt ttg gat gac aac aaa gaa aaa gat gtt agt aag aaa aga aaa      144
248 Tyr Phe Leu Asp Asp Asn Lys Glu Lys Asp Val Ser Lys Lys Arg Lys
249          35          40          45
251 gct caa ttg aag gat gaa tat gat aat ata tca agg agc aaa gaa aat      192
252 Ala Gln Leu Lys Asp Glu Tyr Asp Asn Ile Ser Arg Ser Lys Glu Asn
253          50          55          60
255 att aat aat tcc aaa aaa ata aaa aat gaa tta agt ata aaa gat aat      240
256 Ile Asn Asn Ser Lys Lys Ile Lys Asn Glu Leu Ser Ile Lys Asp Asn
257 65          70          75          80
259 atg cac gat tac ata tat gat gat cgt atc tac aat aat gat aaa gag      288
260 Met His Asp Tyr Ile Tyr Asp Asp Arg Ile Tyr Asn Asn Asp Lys Glu
261          85          90          95
263 aaa aat gtt ata aaa agt gat aat aaa aat gtt ata aaa agt gat aat      336
264 Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys Ser Asp Asn
265          100          105          110
267 aaa aat gat tat aaa aag tgt aat aaa aat gtt ata aaa agt gat aat      384
268 Lys Asn Asp Tyr Lys Lys Cys Asn Lys Asn Val Ile Lys Ser Asp Asn
269          115          120          125
271 aaa aat gtt ata aaa agt gat aat aaa aat gtt ata aaa agt gat aat      432
272 Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys Ser Asp Asn
273          130          135          140
275 aaa aat gtt ata aaa agt gat tat aaa agt gat gat aga aat gct tgt      480
276 Lys Asn Val Ile Lys Ser Asp Tyr Lys Ser Asp Asp Arg Asn Ala Cys
277 145          150          155          160
279 gat att tat aaa agt aat aaa aaa aat gtt cct gat aat tgc cat ata      528
280 Asp Ile Tyr Lys Ser Asn Lys Lys Asn Val Pro Asp Asn Cys His Ile
281          165          170          175
283 tat gat gat aat agt tca gtt gaa aat tta gat gga aaa aat aaa tta      576
284 Tyr Asp Asp Asn Ser Ser Val Glu Asn Leu Asp Gly Lys Asn Lys Leu
285          180          185          190
287 aat aat ata agg aac ata cat aat gat aac tca tct tca tgc gat ata      624
288 Asn Asn Ile Arg Asn Ile His Asn Asp Asn Ser Ser Ser Cys Asp Ile

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| | | | | |
|-----|---|------|-----|--|
| 289 | 195 | 200 | 205 | |
| 291 | tcc gat ata aaa agt gaa gat gaa tat ata gaa cca tat gaa aaa aag | 672 | | |
| 292 | Ser Asp Ile Lys Ser Glu Asp Glu Tyr Ile Glu Pro Tyr Glu Lys Lys | | | |
| 293 | 210 215 220 | | | |
| 295 | aat gaa gaa aat ata aat gaa tat aag aat aag aaa aat ata gcc aat | 720 | | |
| 296 | Asn Glu Glu Asn Ile Asn Glu Tyr Lys Asn Lys Lys Asn Ile Ala Asn | | | |
| 297 | 225 230 235 240 | | | |
| 299 | gaa aat ata aaa gaa gga aag agt tca att tat aat gat gaa cat aat | 768 | | |
| 300 | Glu Asn Ile Lys Glu Gly Lys Ser Ser Ile Tyr Asn Asp Glu His Asn | | | |
| 301 | 245 250 255 | | | |
| 303 | tat aat tca tta tta tat aat tct tgt aat ggt gaa ata agt aag atc | 816 | | |
| 304 | Tyr Asn Ser Leu Leu Tyr Asn Ser Cys Asn Gly Glu Ile Ser Lys Ile | | | |
| 305 | 260 265 270 | | | |
| 307 | aac aaa ata agt agt cat aat aat att gat aat aat atg gat aat tat | 864 | | |
| 308 | Asn Lys Ile Ser Ser His Asn Asn Ile Asp Asn Asn Met Asp Asn Tyr | | | |
| 309 | 275 280 285 | | | |
| 311 | aat acg ttt gca aat gtg aat aat ttt ata ata tat tcc tca gat gat | 912 | | |
| 312 | Asn Thr Phe Ala Asn Val Asn Asn Phe Ile Ile Tyr Ser Ser Asp Asp | | | |
| 313 | 290 295 300 | | | |
| 315 | gaa gat aat ata tca aat tat tat aat ggt aaa gac gta tta aat gat | 960 | | |
| 316 | Glu Asp Asn Ile Ser Asn Tyr Tyr Asn Gly Lys Asp Val Leu Asn Asp | | | |
| 317 | 305 310 315 320 | | | |
| 319 | gag att atg ttc cct ata aaa ttt aat ttt gaa aaa tta aaa aaa aat | 1008 | | |
| 320 | Glu Ile Met Phe Pro Ile Lys Phe Asn Phe Glu Lys Leu Lys Lys Asn | | | |
| 321 | 325 330 335 | | | |
| 323 | att tat gta ata gag cat ata gac aaa ata tat tat gat aca ttt tta | 1056 | | |
| 324 | Ile Tyr Val Ile Glu His Ile Asp Lys Ile Tyr Tyr Asp Thr Phe Leu | | | |
| 325 | 340 345 350 | | | |
| 327 | aat aaa aat cca agt gaa aaa agt gtt ttt atg aat gat gaa tct act | 1104 | | |
| 328 | Asn Lys Asn Pro Ser Glu Lys Ser Val Phe Met Asn Asp Glu Ser Thr | | | |
| 329 | 355 360 365 | | | |
| 331 | ggt tat ttg aaa aat gat gtg aat gac aaa tgt gtt gtt gat aat ata | 1152 | | |
| 332 | Gly Tyr Leu Lys Asn Asp Val Asn Asp Lys Cys Val Val Asp Asn Ile | | | |
| 333 | 370 375 380 | | | |
| 335 | aat gtt att aat cct tct agt gtg aat acg ttg agt aat att tca aat | 1200 | | |
| 336 | Asn Val Ile Asn Pro Ser Ser Val Asn Thr Leu Ser Asn Ile Ser Asn | | | |
| 337 | 385 390 395 400 | | | |
| 339 | att agg aat gaa aaa ata gaa aat aat aat aag aat gaa aaa tta ata | 1248 | | |
| 340 | Ile Arg Asn Glu Lys Ile Glu Asn Asn Asn Lys Asn Glu Lys Leu Ile | | | |
| 341 | 405 410 415 | | | |
| 343 | aaa tca tat cct aca caa tca aaa aat gtt atg agt aca ttt tcc ttt | 1296 | | |
| 344 | Lys Ser Tyr Pro Thr Gln Ser Lys Asn Val Met Ser Thr Phe Ser Phe | | | |
| 345 | 420 425 430 | | | |
| 347 | tgg aat att gaa aag gag aca ttt ata aca aaa cct ttg tat gca caa | 1344 | | |
| 348 | Trp Asn Ile Glu Lys Glu Thr Phe Ile Thr Lys Pro Leu Tyr Ala Gln | | | |
| 349 | 435 440 445 | | | |
| 351 | aat ttg aga aaa aaa caa ttt agt tta tta gat gaa tct gaa gag atg | 1392 | | |
| 352 | Asn Leu Arg Lys Lys Gln Phe Ser Leu Leu Asp Glu Ser Glu Glu Met | | | |
| 353 | 450 455 460 | | | |

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355 ata aga aat tat tca tct aat caa tat tct ata aaa ttt gta cca aga      1440
356 Ile Arg Asn Tyr Ser Ser Asn Gln Tyr Ser Ile Lys Phe Val Pro Arg
357 465                      470                      475                      480
359 cat tta tta tat gta atg agt caa gtt gct tct cga tcc ttt ttt gat      1488
360 His Leu Leu Tyr Val Met Ser Gln Val Ala Ser Arg Ser Phe Phe Asp
361                      485                      490                      495
363 cct tta tat aga aag cag tta ttt ttt cgt tac taa      1524
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368 <210> SEQ ID NO: 10
370 <211> LENGTH: 507
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374 <213> ORGANISM: Plasmodium falciparum
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385                      20                      25                      30
388 Tyr Phe Leu Asp Asp Asn Lys Glu Lys Asp Val Ser Lys Lys Arg Lys
389                      35                      40                      45
392 Ala Gln Leu Lys Asp Glu Tyr Asp Asn Ile Ser Arg Ser Lys Glu Asn
393                      50                      55                      60
396 Ile Asn Asn Ser Lys Lys Ile Lys Asn Glu Leu Ser Ile Lys Asp Asn
397 65                      70                      75                      80
400 Met His Asp Tyr Ile Tyr Asp Asp Arg Ile Tyr Asn Asn Asp Lys Glu
401                      85                      90                      95
404 Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys Ser Asp Asn
405                      100                     105                     110
408 Lys Asn Asp Tyr Lys Lys Cys Asn Lys Asn Val Ile Lys Ser Asp Asn
409                      115                     120                     125
412 Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys Ser Asp Asn
413                      130                     135                     140
416 Lys Asn Val Ile Lys Ser Asp Tyr Lys Ser Asp Asp Arg Asn Ala Cys
417 145                     150                     155                     160
420 Asp Ile Tyr Lys Ser Asn Lys Lys Asn Val Pro Asp Asn Cys His Ile
421                      165                     170                     175
424 Tyr Asp Asp Asn Ser Ser Val Glu Asn Leu Asp Gly Lys Asn Lys Leu
425                      180                     185                     190
428 Asn Asn Ile Arg Asn Ile His Asn Asp Asn Ser Ser Ser Cys Asp Ile
429                      195                     200                     205
432 Ser Asp Ile Lys Ser Glu Asp Glu Tyr Ile Glu Pro Tyr Glu Lys Lys
433                      210                     215                     220
436 Asn Glu Glu Asn Ile Asn Glu Tyr Lys Asn Lys Lys Asn Ile Ala Asn
437 225                     230                     235                     240
440 Glu Asn Ile Lys Glu Gly Lys Ser Ser Ile Tyr Asn Asp Glu His Asn
441                      245                     250                     255
444 Tyr Asn Ser Leu Leu Tyr Asn Ser Cys Asn Gly Glu Ile Ser Lys Ile
445                      260                     265                     270
448 Asn Lys Ile Ser Ser His Asn Asn Ile Asp Asn Asn Met Asp Asn Tyr

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VERIFICATION SUMMARY

DATE: 08/10/2004

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TIME: 11:50:30

Input Set : A:\PTO.LM.txt

Output Set: N:\CRF4\08102004\J712533.raw

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L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:238 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:9,Line#:234
L:538 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:11,Line#:524
L:548 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:11,Line#:534
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